Amendments to the Claims

The listing of claims will replace all prior versions, and listings of claims in the application.

1-19. (cancelled).

- 20. (currently amended) A method of screening for an agonist or an antagonist of PTH receptor activity comprising:
- (a) contacting cells with a test compound wherein said cells express a $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having an amino acid sequence at least 95% identical to a sequence selected from the group consisting of:
- (i) the amino acid sequence from about position 1 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (ii) the amino acid sequence from about position 2 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iii) the amino acid sequence from about position 23 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iv) the amino acid sequence of the $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No.

PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted; and

(v) the amino acid sequence of the mature <u>rΔNt</u> [[rδNt]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted;

wherein said polypeptide increases intracellular cAMP levels when activated by PTH or PTH-related peptide extracellular amino-terminal ligand binding domain has an amino acid sequence from about residue 26 to about residue 181 in wild-type PTH receptor;

- (b) measuring cAMP accumulation in said cells; and
- (c) determining whether said test compound is an agonist or an antagonist of PTH receptor activity:

wherein an agonist is identified as a compound that increases cAMP accumulation and an antagonist prevents cAMP accumulation.

- 21. (cancelled).
- 22. (previously presented) The method of claim 20, wherein said agonist is a peptide.
- 23. (previously presented) The method of claim 20, wherein said antagonist is a peptide.

- 24. (currently amended) A method of screening for an agonist or an antagonist of PTH receptor activity comprising:
- (a) contacting cells with a test compound wherein said cells express a $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide, wherein said cells comprise a polynucleotide having a nucleotide sequence at least 95% identical to a sequence selected from the group consisting of:
- (i) a nucleotide sequence encoding the amino acid sequence from about position 1 to about position 1320 in SEQ ID NO:1-435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (ii) a nucleotide sequence encoding the amino acid sequence from about position 4 to about position 1320 in SEQ ID NO:1-2 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iii) a nucleotide sequence encoding the amino acid sequence from about position 67 to about position 1320 in SEQ ID NO:1 23 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iv) a nucleotide sequence encoding the $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted; and
- (v) a nucleotide sequence encoding the mature $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in

ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted;

wherein said polypeptide increases intracellular cAMP levels when activated by PTH or PTH-related peptide extracellular amino-terminal ligand binding domain has an amino acid sequence from about residue 26 to about residue 181 in wild-type PTH receptor;

- (b) measuring cAMP accumulation in said cells; and
- (c) determining whether said test compound is an agonist or an antagonist of PTH receptor activity;

wherein an agonist is identified as a compound that increases cAMP accumulation and an antagonist prevents cAMP accumulation.

- 25. (currently amended) A method of screening for an agonist or an antagonist of PTH receptor activity comprising:
- (a) contacting cells with a test compound wherein said cells express a $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having an amino acid sequence selected from the group consisting of:
- (i) the amino acid sequence from about position 1 to about position 435 in SEQ ID NO:2;
- (ii) the amino acid sequence from about position 2 to about position 435 in SEQ ID NO:2;
- (iii) the amino acid sequence from about position 23 to about position 435 in SEQ ID NO:2;

- (iv) the amino acid sequence of the $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136; and
- (v) the amino acid sequence of the mature $\underline{r\Delta Nt}$ [[r δNt]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136;

wherein said polypeptide comprises a deletion of the extracellular amino-terminal ligand binding domain of a PTH-1 receptor, said extracellular amino-terminal ligand binding domain having an amino acid sequence from about residue 26 to about residue 181 in wild-type PTH receptor;

- (b) measuring cAMP accumulation in said cells; and
- (c) determining whether said test compound is an agonist or an antagonist of PTH receptor activity;

wherein an agonist is identified as a compound that increases cAMP accumulation and an antagonist prevents cAMP accumulation.

- 26. (currently amended) A method of screening for an agonist or an antagonist of PTH receptor activity comprising:
- (a) contacting cells with a test compound wherein said cells express a <u>rΔNt</u>
 [[rδNt]] polypeptide, wherein said cells comprise a polynucleotide having a nucleotide sequence selected from the group consisting of:
- (i) a nucleotide sequence encoding the amino acid sequence from about position 1 to about position 435 in SEQ ID NO:2;

- (ii) a nucleotide sequence encoding the amino acid sequence from about position 2 to about position 435 in SEQ ID NO:2;
- (iii) a nucleotide sequence encoding the amino acid sequence from about position 23 to about position 435 in SEQ ID NO:2;
- (iv) a nucleotide sequence encoding the $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136; and
- (v) a nucleotide sequence encoding of the mature $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136;

wherein said polypeptide comprises a deletion of the extracellular amino-terminal ligand binding domain of a PTH-1 receptor, said extracellular amino-terminal ligand binding domain having an amino acid sequence from about residue 26 to about residue 181 in wild-type PTH receptor;

- (b) measuring the biological response of cAMP accumulation in said cells;
- (c) determining whether said test compound is an agonist or an antagonist of PTH receptor activity;

 wherein an agonist is identified as a compound that increases cAMP accumulation and an antagonist prevents cAMP accumulation.
- 27. (currently amended) A method of screening for an agonist or an antagonist of PTH receptor activity comprising:

- (a) providing an iodinated test compound;
- (b) contacting cells with said iodinated test compound wherein said cells express a $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide, wherein said cells comprise a polynucleotide having a nucleotide sequence at least 95% identical to a sequence selected from the group consisting of:
- (i) a nucleotide sequence encoding the amino acid sequence from about position 1 to about position 1320 in SEQ ID NO:1-435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (ii) a nucleotide sequence encoding the amino acid sequence from about position 4 to about position 1320 in SEQ ID NO:1-2 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iii) a nucleotide sequence encoding the amino acid sequence from about position 67 to about position 1320 in SEQ ID NO:1 23 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iv) a nucleotide sequence encoding the $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted; and
- (v) a nucleotide sequence encoding the mature $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted;

wherein said polypeptide increases intracellular cAMP levels when activated by PTH or PTH-related peptide extracellular amino-terminal ligand binding domain has an amino acid sequence from about residue 26 to about residue 181 in wild-type PTH receptor; and

- (b) determining whether said iodinated test compound competitively binds to said <u>rΔNt</u> [[rδNt]] polypeptide; wherein an agonist is identified as a compound that increases cAMP accumulation and an antagonist prevents cAMP accumulation.
- 28. (currently amended) A method of screening for an agonist or an antagonist of PTH receptor activity comprising:
 - (a) providing an iodinated test compound;
- (b) contacting cells with said iodinated test compound wherein said cells express a $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having an amino acid sequence at least 95% identical to a sequence selected from the group consisting of:
- (i) the amino acid sequence from about position 1 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (ii) the amino acid sequence from about position 2 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;

- (iii) the amino acid sequence from about position 23 to about position 435 in SEQ ID NO:2, wherein the extracellular amino-terminal ligand binding domain is deleted;
- (iv) the amino acid sequence of the $\underline{r\Delta Nt}$ [[$r\delta Nt$]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted; and
- (v) the amino acid sequence of the mature $\underline{r\Delta Nt}$ [[r δNt]] polypeptide having the amino acid sequence encoded by the cDNA clone contained in ATCC Deposit No. PTA-1136, wherein the extracellular amino-terminal ligand binding domain is deleted;

wherein said polypeptide increases intracellular cAMP levels when activated by PTH or PTH-related peptide extracellular amino-terminal ligand binding domain has an amino acid sequence from about residue 26 to about residue 181 in wild-type PTH receptor; and

(b) determining whether said iodinated test compound competitively binds to said <u>rΔNt</u> [[rδNt]] polypeptide; wherein an agonist is identified as a compound that increases cAMP accumulation and an antagonist prevents cAMP accumulation.